

F i g . 1

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C A K S P L E N T W W A E Q D S P F C F L A F C F E :517
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S Q L D K M P A L P A K G N L N L R D I L E S D F A :881
TTCCGGT:5820
F A :883

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T7 436:GNDMTKGLTLAKGKPI-GKEGYWLKIHGANCAGVDKVFFPERI-KFIEENHENI-MACAKSPLENTWAEQDSPF
T3 437:.....E..F.....-A...KHVDD.-L....D.IN.....
K11 459:.....S.....LD.F.....EG...-L.S.AD.N....TQ.....
SP6 427:S..LG.A..RFTE.R.VN.V.ALK.FC.N...LW.W..KT.DV.VSNULD.EFQDMCRDI.AD..TF.Q...KA.A.Y
** *

T7 513:AFCFEYA---GVQHGG-L-SYNCSLPLAFDGCSCGIQHFSAMLRDEVGGRAVNLLPSETVQDIYGIVAKVNILQA
T3 514:.....T.....
K11 536:.....K.....-N.....SI.....D.....K..D....V.HQ
SP6 507:W....QYLDLVDE.RADEFRTH..VHQ.....Y.....AK....K.DAP.....A.-Q.--V-I
* **** * ** ***** * **** * * **** * * * *

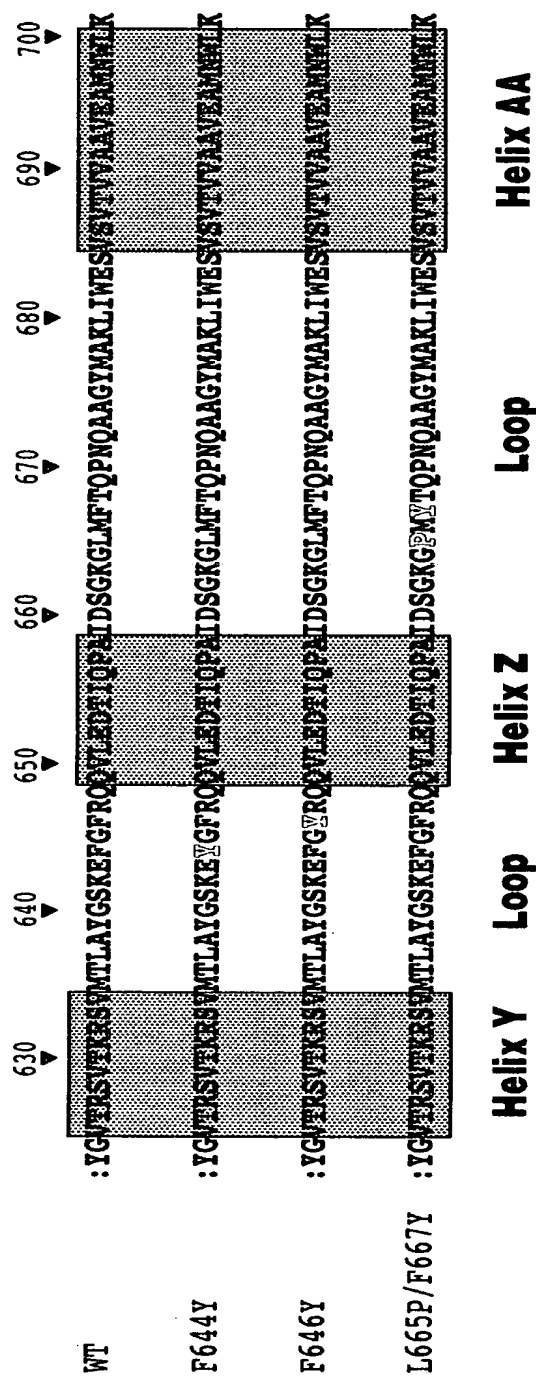
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T3 589:..P..MI....KD.....L....ST..Q.....D.....
K11 611:..SQTVEQIA.KE..FH..T..ESV..A..Q....K.....SLV.....N.E
SP6 581:ALYMDADDA.TFTS.SVTLSGT-ELR.M.SA.DSI.I..L..KP....P...TRLTC.ES.IDYIVDLEEKEAQ.
* *

T7 667:FT-QPNQAAGYMAKLWESVSVTVAAVEAMNLWSAAKLAAEVKDKKTGEILR---KRC-AVHWVTPDGFPVWQE
T3 668:..-.....DA.....K.....K.V-----H...T.....
K11 690:..H.....DA.T.....K.....K.V-----I.....
SP6 660:EGRTA.KVHPFEDDRQDYLTPGAANYMT.LI.PSISEVVK.PI.AM.MIROLA.FAA.NEGLMYTL.T..ILE.K
* *

T7 742:PIQTRNLNMFGLQFLQPTINTNKDSEIDAHAHQESGIAPNFVHSQDSHLRKTVVWAHEKYGIESFALIHDSSFGTIP
T3 743:L.K.DMI.....L..G.....M...Y.....
K11 765:QN.A..K.V...ANVKM.Y.G.....ANVKM.Y.G.....M...H.N.V..D.....S....
SP6 740:TEML.VRTCLM.DIKMSLOVE.--IV.EAAMMGAA.....GH.A...IL.CELVD--VT.I.V.....HA
* *

T7 822:ANLFKAVRETMVDTYESCDVLADFYDQFADQLHESOLDKMPALPAKNLNLRDILESDFABA
T3 823:GK.....I..NN.....T.....P.K....Q..K.....
K11 845:G.....K..DN..I.....V.....D.....
SP6 816:LT.RV.LKGQ..AM.IDGNA.QKL-LE-EHEVR-WMV.TGIEV.EQ.EFD.NE.MD.EVV..
* *

Fig. 5



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[illegible]

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T7RNaPol 420:MDWRGRVYAVSMFNPQGNMTKGLLTLAKGKPIGEGYYWLKIHGANCAGVDKVPFFPERI
T3RNaPol 421:.....P.....E..F.....
*****
T7RNaPol 480:KFIEENHENIMACAKSPLENTWAEQDSPFCFLAFCFAYAGVQHGLSYNCSLPLAFDGS
T3RNaPol 481:A...KHVDD.L...D.IN.....T.....
*****
T7RNaPol 540:CSGIGHFSAMLRDEVGGRAVNLLPSETVQDIYIVAKKVNEILOADAINGTDNEVTVTVD
T3RNaPol 541:.....Q.....KQ.....P..MI.....
*****
T7RNaPol 600:ENTGEISEKVKLGTKALAGQWLAYGVTRSVTKRSVMTLAYGSKFEFRQVLEDTIQPAI
T3RNaPol 601:KD.....L...ST..Q.....D.....
*****
T7RNaPol 660:DSGKGLMFTQPNQAAGYMAKLIWESVSVTVVAAVEAMNWLKSAKLLAEVVDKKTGEIL
T3RNaPol 661:.....DA.....K...
*****
T7RNaPol 720:RKRCVHVWVTPDGFVPWQOEYKPKIQTRLNLMFLGQFRLQPTINTNKDSEIDAHKQESGIA
T3RNaPol 721:H.....T.....R..L.K.DMI.....L...G.....
*****
T7RNaPol 780:PNFVHSDGSHLRKTVVWAHEKYGIESFALIHDSFGTIPADAANLFKAVRETMVDTYESC
T3RNaPol 781:.....M...Y.....GK.....I...NN
*****
T7RNaPol 840:DVLADFYDQFADQLHESQLDKMPALPAKGNLNLRDILESDFAF
T3RNaPol 841:.....S.....T.....P..K.....Q...K.....
*****

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Fig. 8

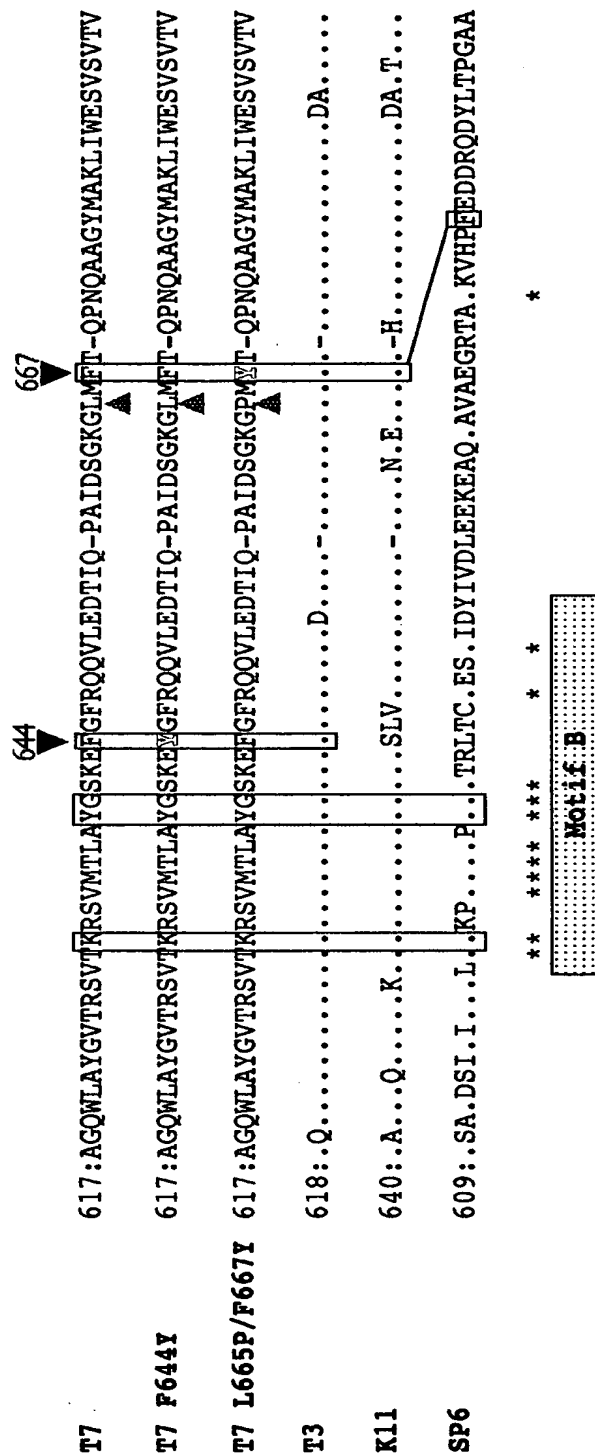


Fig. 9

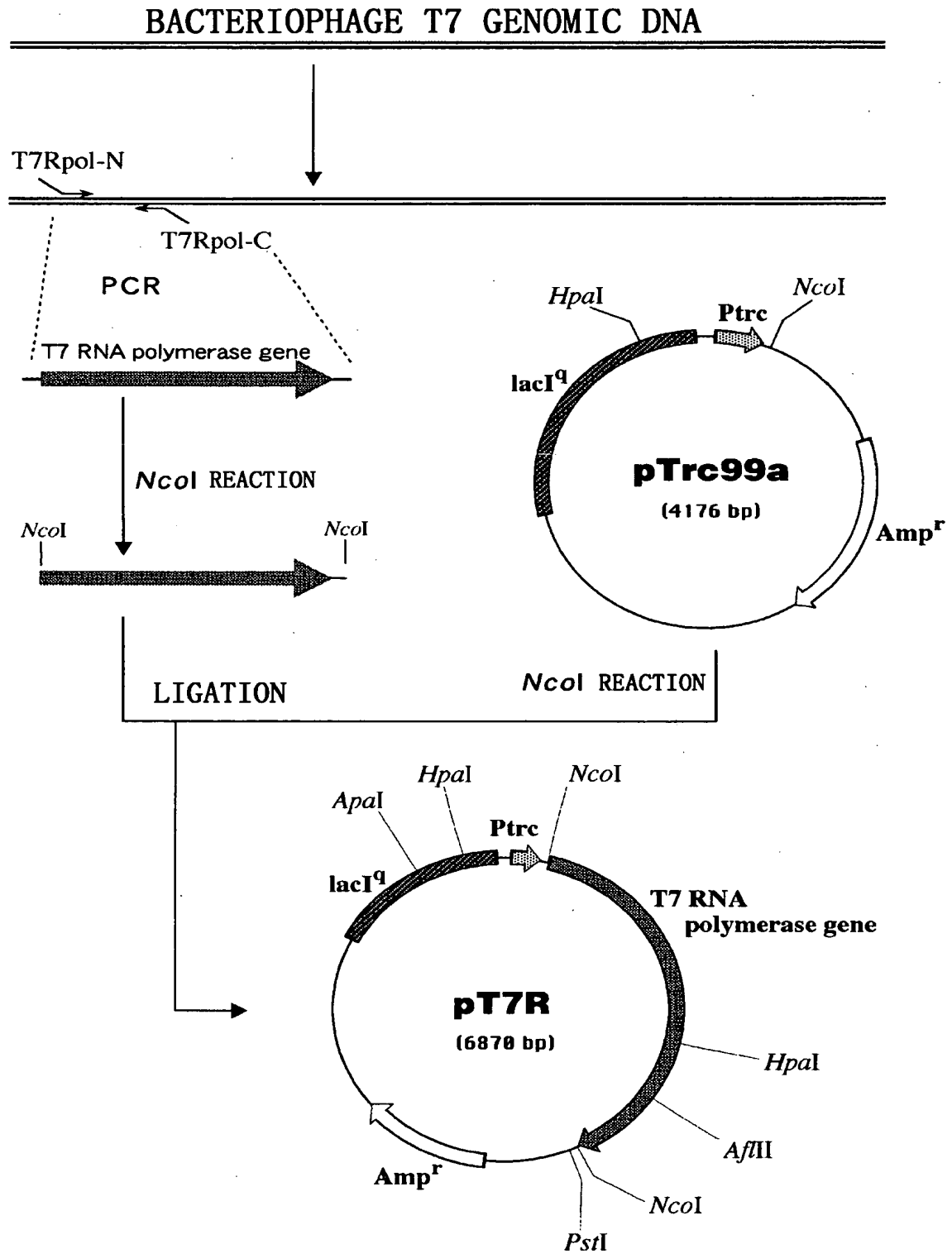


Fig. 10

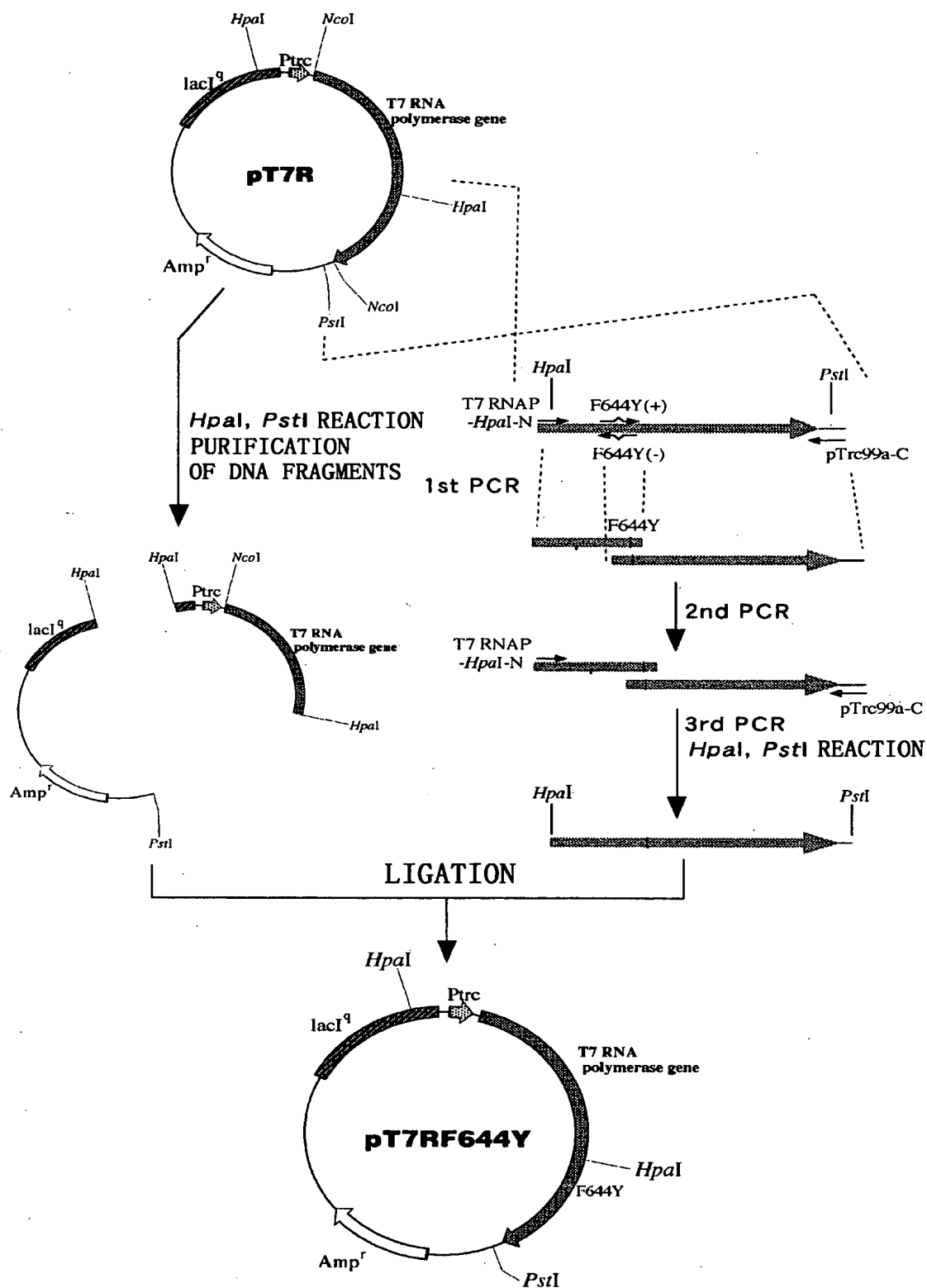


Fig. 11

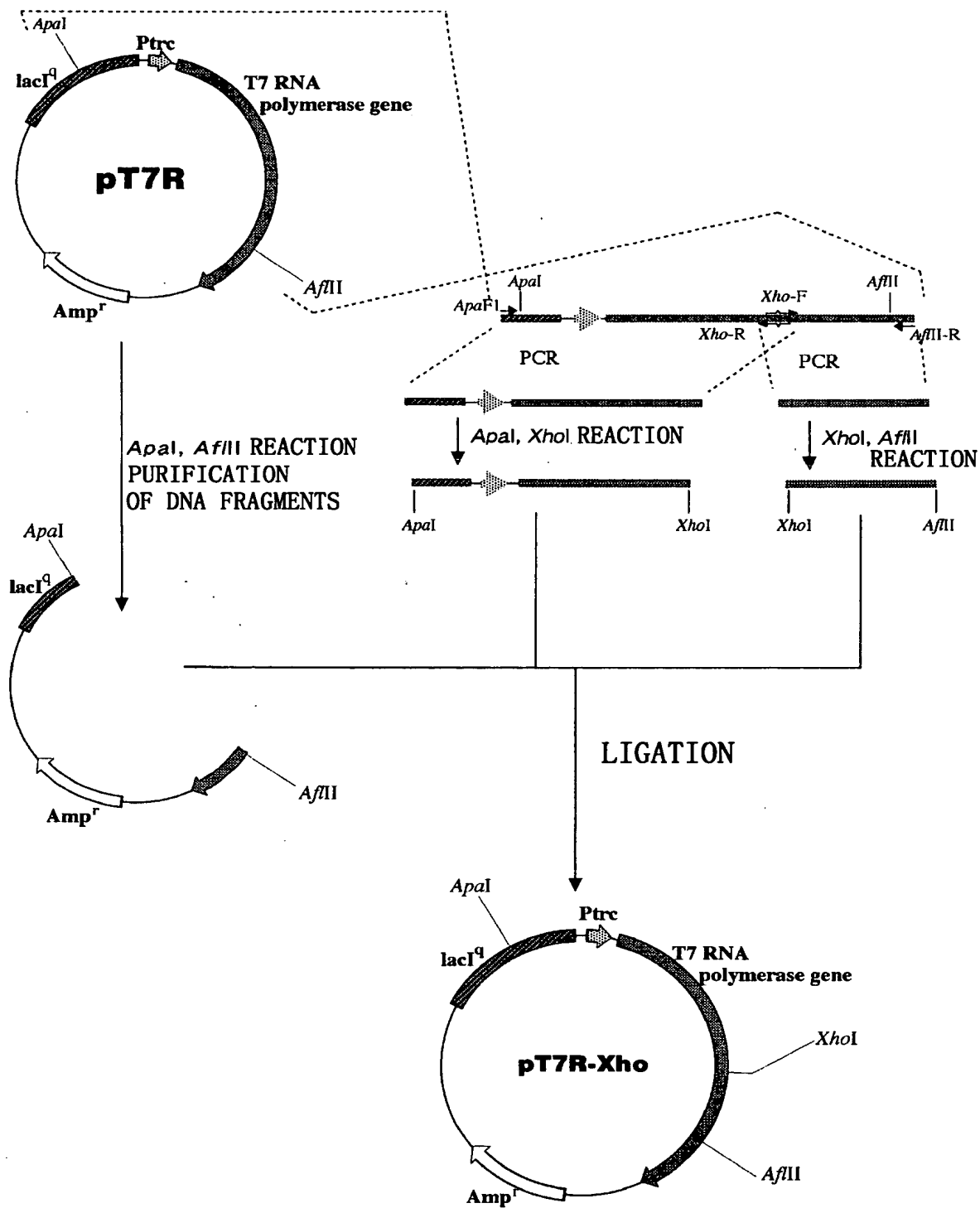


Fig. 12

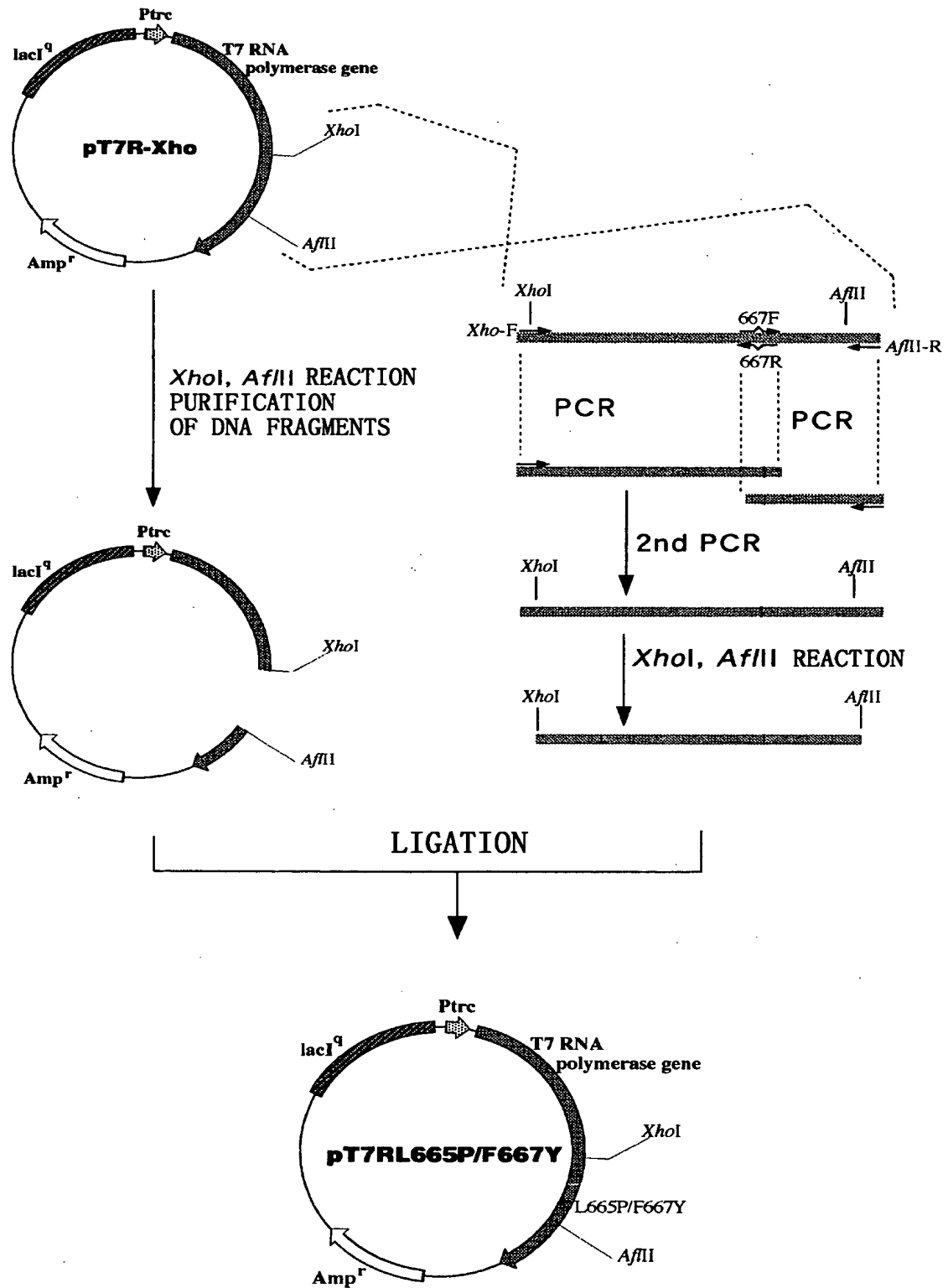


Fig. 13

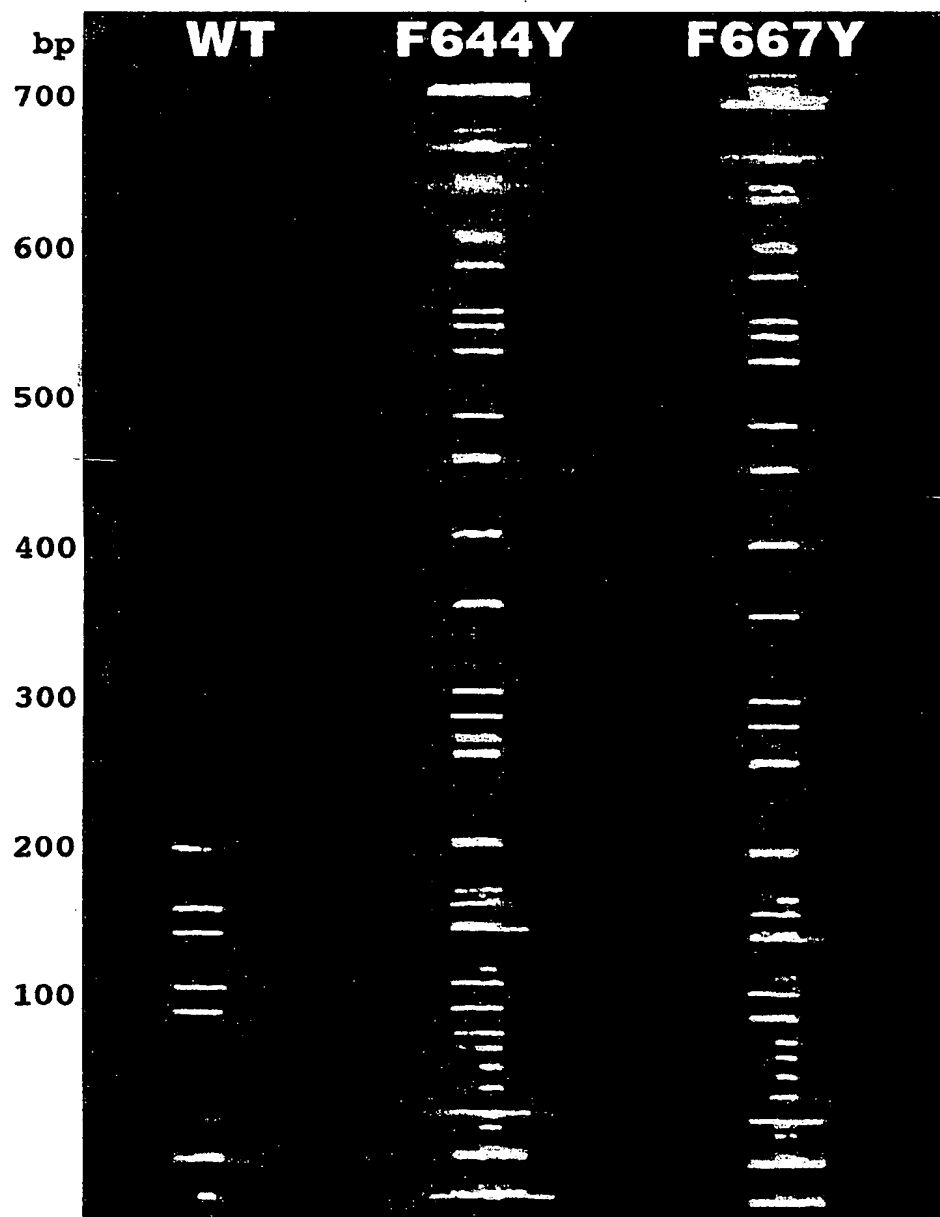


Fig. 14

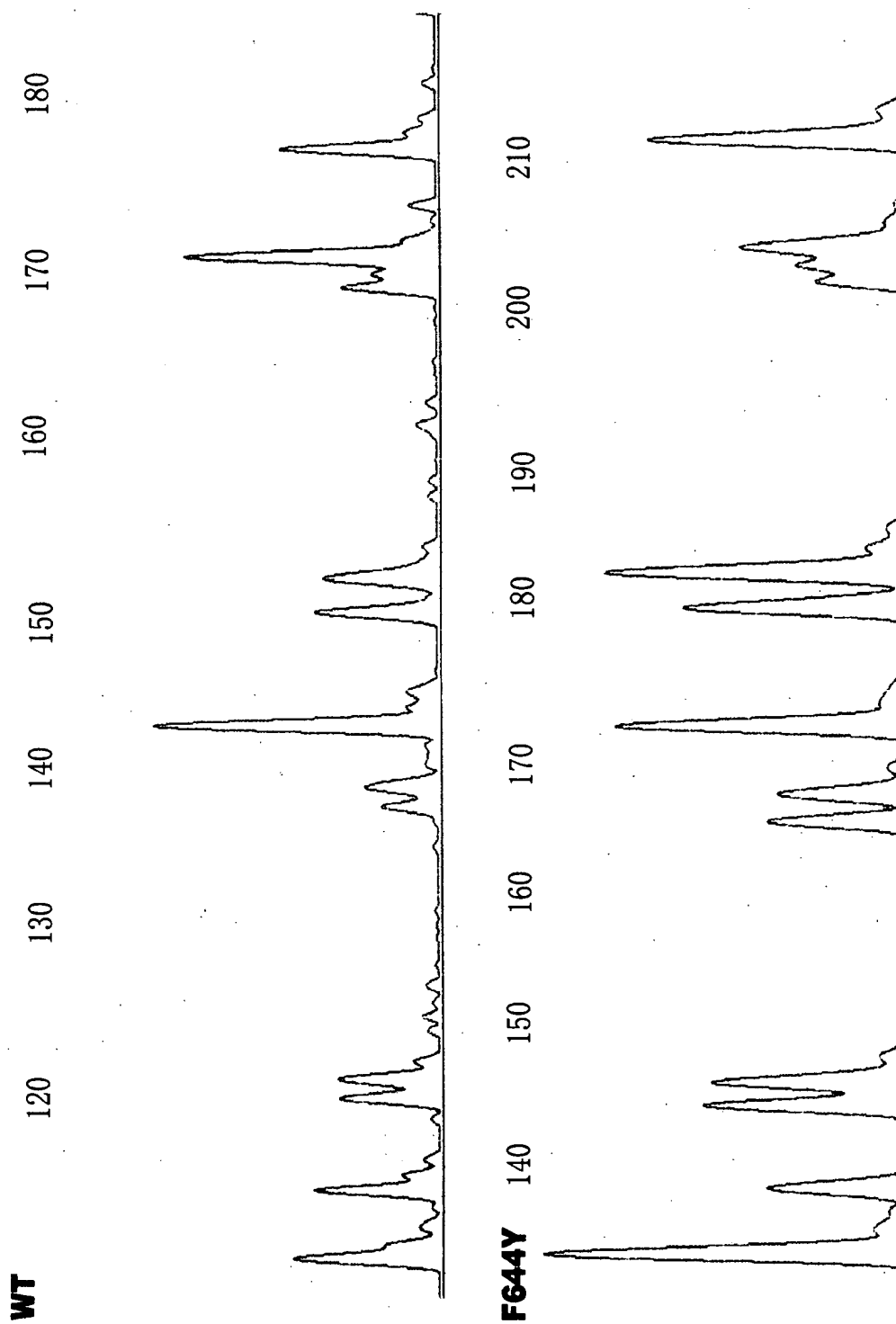


Fig. 15

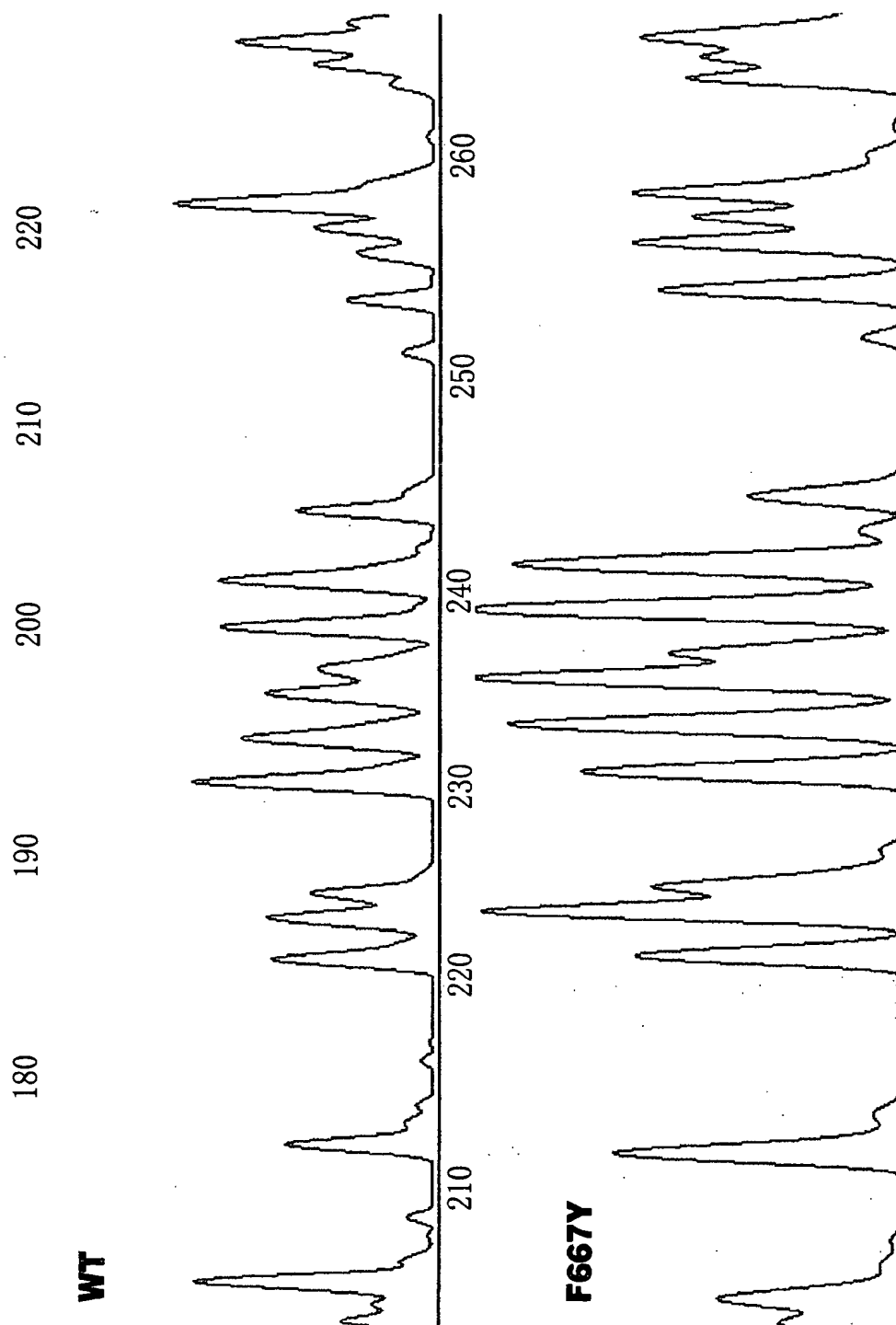


Fig. 16

WT

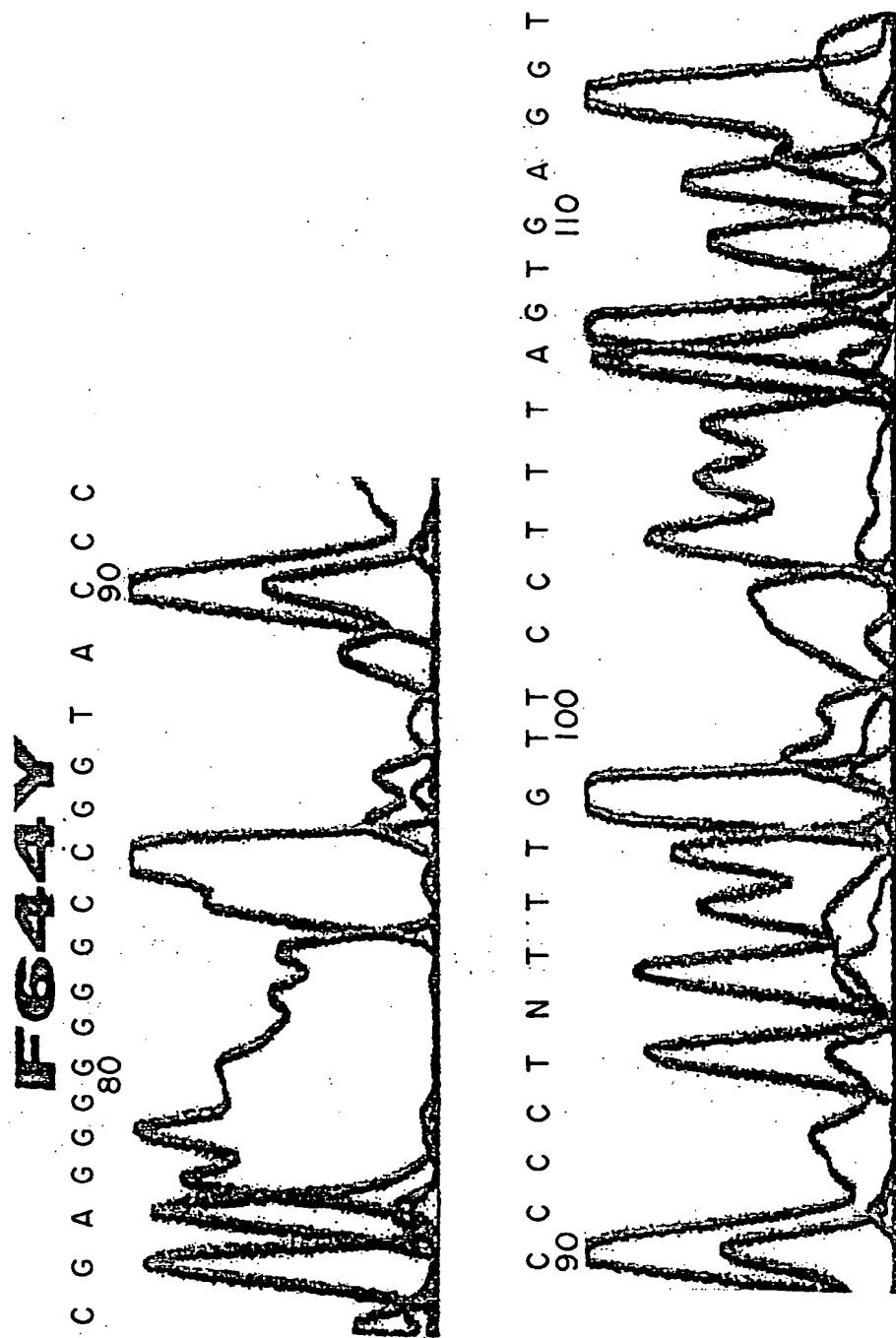
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 200 210 220 230



Fig. 16



1974

S A G G G G G G C C G G T A A C G C
 70 80



A C G C C T T T G T T C C C T T A G T G
 80 90 100

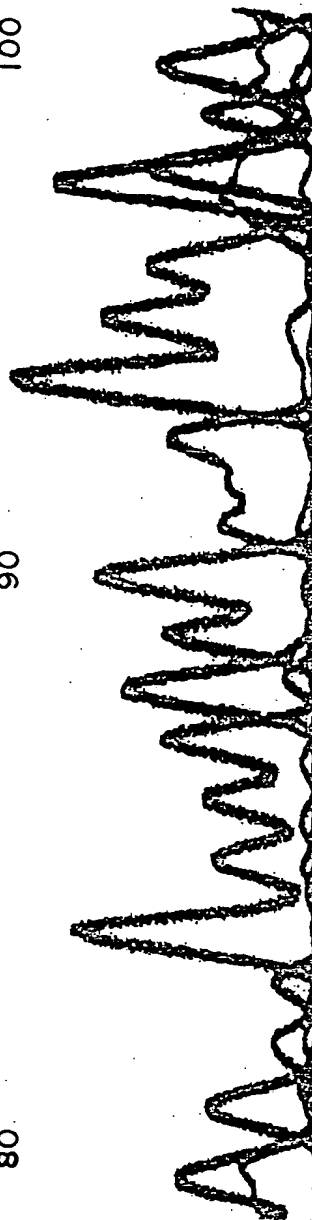


Fig. 17

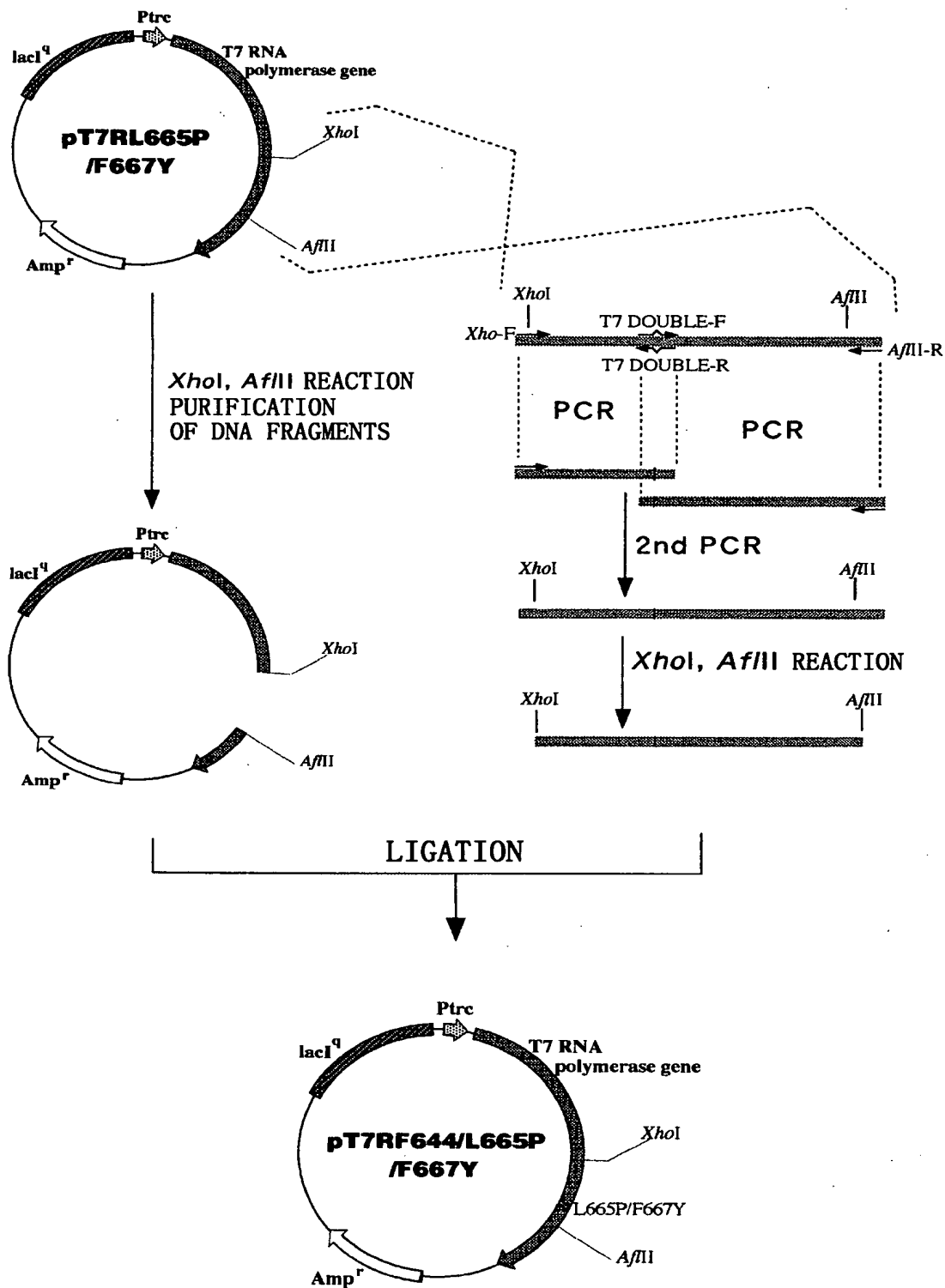


Fig. 18

(1)

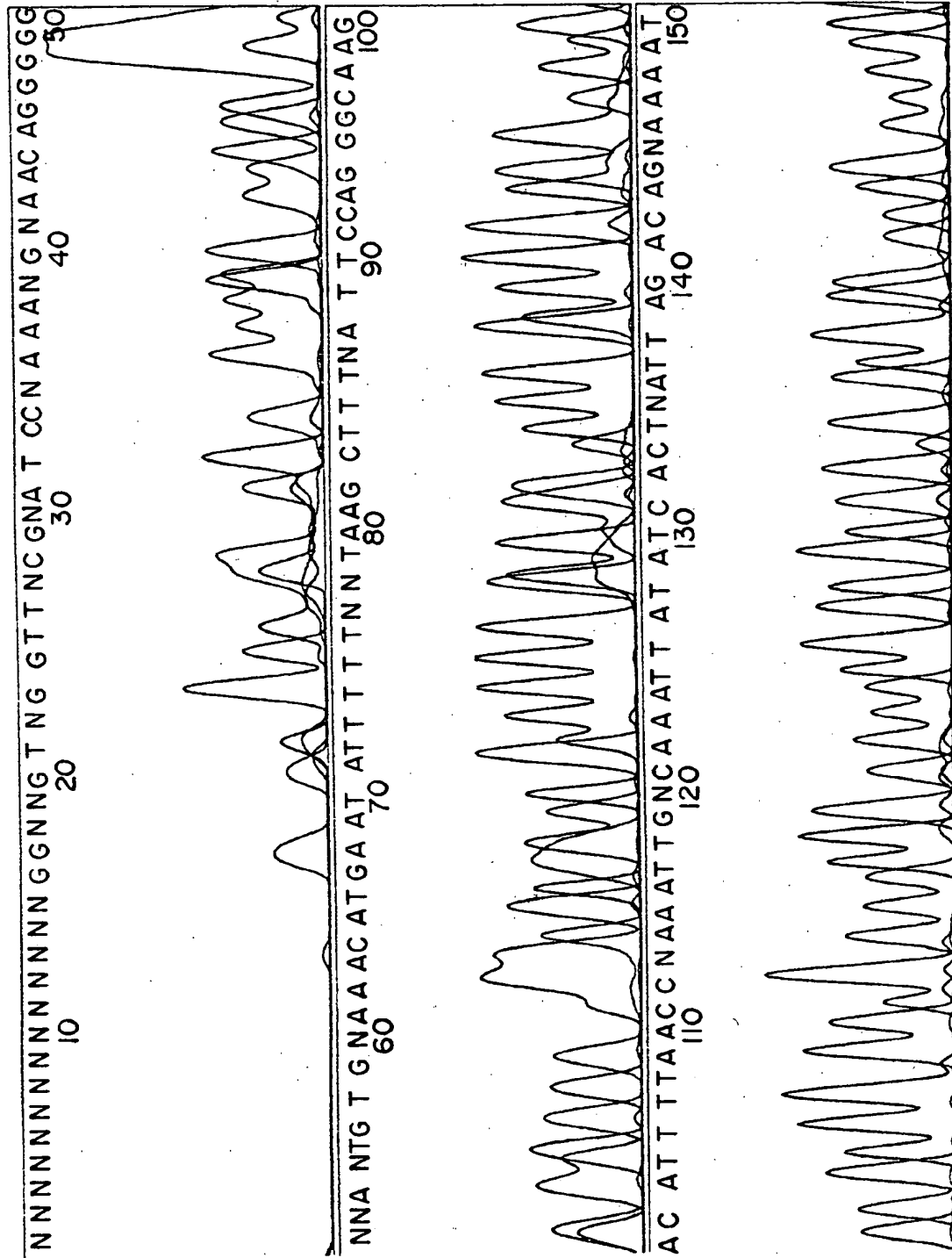


Fig. 18

(2)

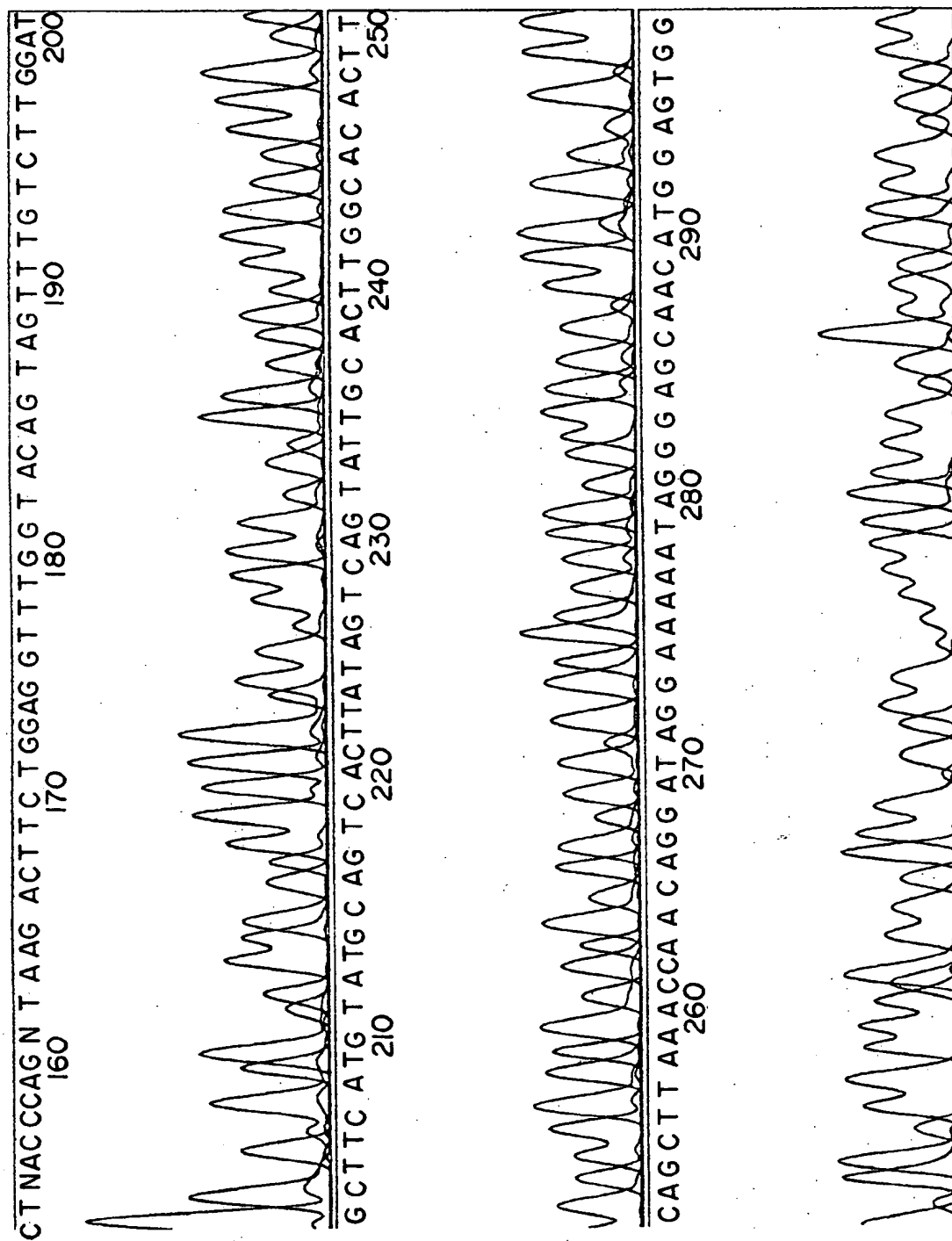


Fig. 18

(3)

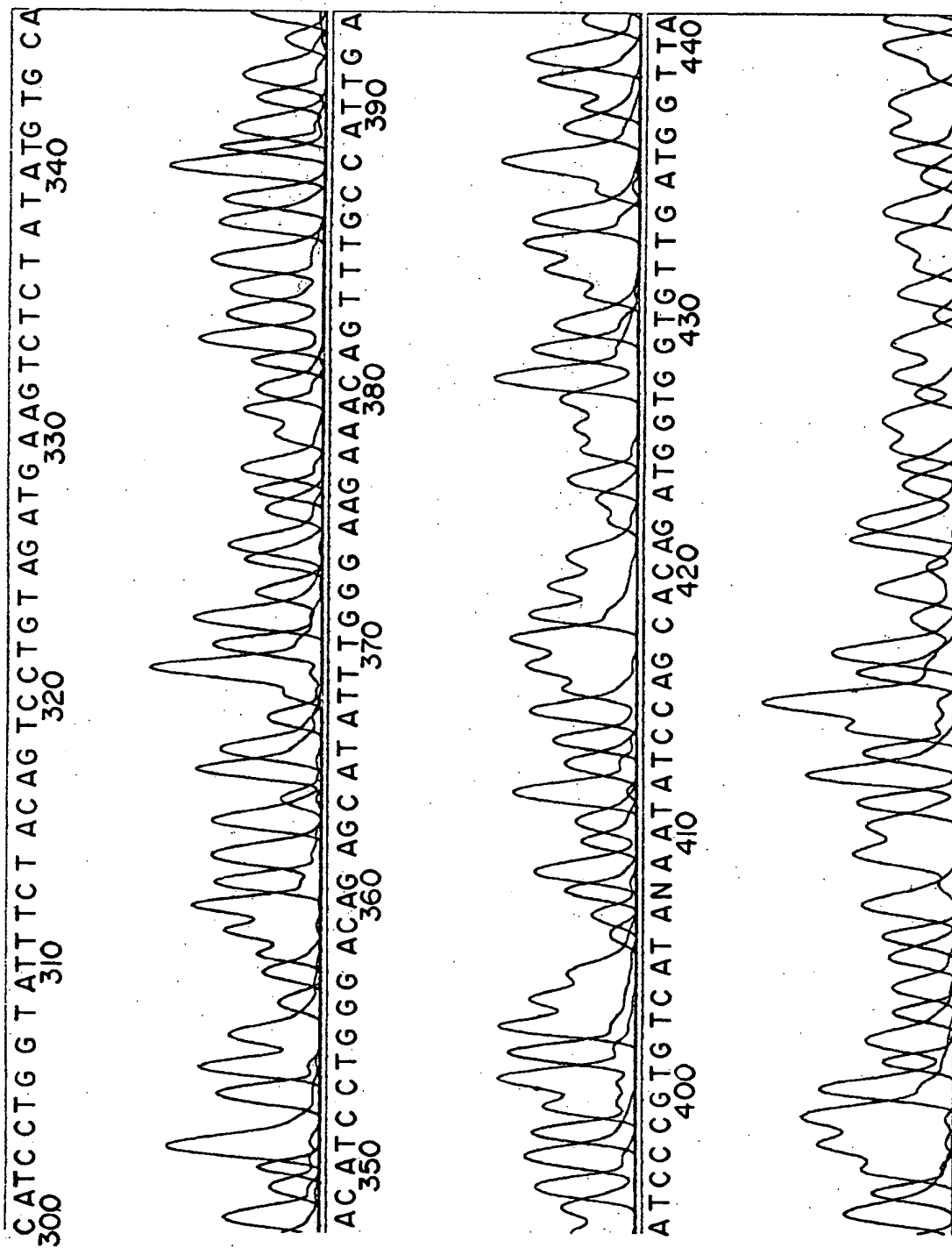


Fig. 18

(4)

